

**IN THE
UNITED STATES
PATENT AND TRADEMARK
OFFICE**

<i>Application Number</i>	09/749,637
<i>Filing Date</i>	28 December 2000
<i>First Named Inventor</i>	Baldomero M. OLIVERA
<i>Group Art Unit</i>	1646
<i>Examiner Name</i>	unassigned
<i>Attorney Docket Number</i>	2314-227

Title of the Invention: **O-SUPERFAMILY CONOTOXIN PEPTIDES**

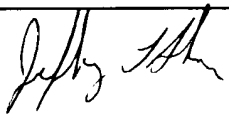
STATEMENT PURSUANT TO 37 CFR 1.821(f)

Assistant Commissioner for Patents
Washington, D.C. 20231

Dear Sir:

In the matter of the above-identified application, Applicants submit herewith a computer disk containing the sequences of the substitute Sequence Listing which is filed concurrently herewith by Preliminary Amendment.

It is hereby certified that the content of the paper and computer copies of said substitute Sequence Listing are identical and include no new matter.

RESPECTFULLY SUBMITTED,					
NAME AND REG. NUMBER	Jeffrey L. Ihnen, Reg. No. 28,957				
SIGNATURE				DATE	13 AUGUST 2001
Address	ROTHWELL, FIGG, ERNST & MANBECK, pc Suite 701-East, 555 13th Street, N.W.				
City	Washington	State	D.C.	Zip Code	20004
Country	U.S.A.	Telephone	202-783-6040	Fax	202-783-6031

RAW SEQUENCE LISTING ERROR REPORT

BIOTECHNOLOGY
SYSTEMS
BRANCH



The Biotechnology Systems Branch of the Scientific and Technical Information Center (STIC) detected errors when processing the following computer readable form:

Application Serial Number: 09/ 749,637

Source: O I P E

Date Processed by STIC: 01-16-01

RECEIVED

AUG 29 2001

TECH CENTER 1600 2900

THE ATTACHED PRINTOUT EXPLAINS DETECTED ERRORS.

PLEASE FORWARD THIS INFORMATION TO THE APPLICANT BY EITHER:

- 1) INCLUDING A COPY OF THIS PRINTOUT IN YOUR NEXT COMMUNICATION TO THE APPLICANT, WITH A NOTICE TO COMPLY or,
- 2) TELEPHONING APPLICANT AND FAXING A COPY OF THIS PRINTOUT, WITH A NOTICE TO COMPLY

FOR CRF SUBMISSION QUESTIONS, PLEASE CONTACT MARK SPENCER, 703-308-4212.

FOR SEQUENCE RULES INTERPRETATION, PLEASE CONTACT ROBERT WAX, 703-308-4216.

PATENTIN 2.1 e-mail help: patin21help@uspto.gov or phone 703-306-4119 (R. Wax)

PATENTIN 3.0 e-mail help: patin30help@uspto.gov or phone 703-306-4119 (R. Wax)

TO REDUCE ERRORED SEQUENCE LISTINGS, PLEASE USE THE CHECKER VERSION 3.0 PROGRAM, ACCESSIBLE THROUGH THE U.S. PATENT AND TRADEMARK OFFICE WEBSITE. SEE BELOW:

Checker Version 3.0

The Checker Version 3.0 application is a state-of-the-art Windows based software program employing a logical and intuitive user-interface to check whether a sequence listing is in compliance with format and content rules. Checker Version 3.0 works for sequence listings generated for the original version of 37 CFR §§1.821 - 1.825 effective October 1, 1990 (old rules) and the revised version (new rules) effective July 1, 1998 as well as World Intellectual Property Organization (WIPO) Standard ST.25.

Checker Version 3.0 replaces the previous DOS-based version of Checker, and is Y2K-compliant. Checker allows public users to check sequence listings in Computer Readable form (CRF) before submitting them to the United States Patent and Trademark Office (USPTO).

Use of Checker prior to filing the sequence listing is expected to result in fewer errored sequence listings, thus saving time and money.

Checker Version 3.0 can be down loaded from the USPTO website at the following address:

<http://www.uspto.gov/web/offices/pac/checker>

Note: The initial sequence listing was errored and edited and is included. The edited sequence listing was found to still be errored as is noted on p. 5.

OIPE

RAW SEQUENCE LISTING

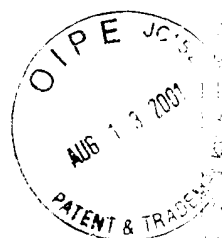
PATENT APPLICATION: US/09/749,637

DATE: 01/23/2001

TIME: 12:19:41

Input Set: A:\Cpg.pto

Output Set: N:\CRF3\01232001\I749637.raw



Does Not Comply
Corrected Diskette Needed
see p. 5

110 APPLICANT: University of Utah Research Foundation
Cognetix, Inc.
Olivera, Baldemero M.
Cartier, G. Edward
Watkins, Marion
Hillyard, David R.
Neimosh, J. Michael
Layor, Richard L.
Jones, Robert M.

115 TITLE OF INVENTION: Superfamily Conserved Peptides

130 FILE REFERENCE: 23 1-227

C--> 33 <140> CURRENT APPLICATION NUMBER: US/09/749,637

C--> 33 <141> CURRENT FILING DATE: 2000-02-28

43 <150> PRIOR APPLICATION NUMBER: US 60/210,412

45 <151> PRIOR FILING DATE: 2000-10-27

46 <150> PRIOR APPLICATION NUMBER: US60/210,440

48 <151> PRIOR FILING DATE: 2000-07-20

49 <150> PRIOR APPLICATION NUMBER: US 60/174,268

47 <151> PRIOR FILING DATE: 2000-06-26

51 <150> PRIOR APPLICATION NUMBER: US 60/173,751

53 <151> PRIOR FILING DATE: 1999-12-30

57 <160> NUMBER OF SEQ ID NOS: 402

61 <170> SOFTWARE: Patent-n Version 3.0

65 <210> SEQ ID NO: 1

67 <211> LENGTH: 261

69 <212> TYPE: DNA

71 <213> ORGANISM: *Comus gloriamaris*

75 <220> FEATURE:

77 <221> NAME/KEY: CDS

79 <222> LOCATION: (1)..(231)

83 <100> SEQUENCE: 1

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86 1          5          10          15
88 tgg acn ttc gtc acg gct gat gac tcc gga aat gga atg qaq att ctt      96
89 Trp Thr Phe Val Thr Ala Asp Asp Ser Gly Asn Gly Met Glu Ile Leu
90          20          25          30
92 ttt ccc aag gcc ggt cac gaa atg qaq aac ctc gaa gtc tct aat ccg      144
93 Phe Pro Lys Ala Gly His Glu Met Glu Asn Leu Glu Val Ser Asn Arg
94          35          40          45
96 gtc aag ccg tgc cgt aaa gaa ggt caa ctt tct gat ccg ata ttt caa      192
97 Val Lys Pro Cys Arg Lys Gly Gln Leu Cys Asp Pro Ile Phe Gln
98          50          55          60
100 aac tgc tgc cgt gcc tgg aat tgc gtt ctt ttc tgc gtc tgaactacc      241
101 Asn Cys Cys Arg Gly Trp Asn Cys Val Leu Phe Cys Val
102 65          70          75
104 gtgatgtctt ctctccctc      261

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RAW SEQUENCE LISTING

PATENT APPLICATION: US/09/749,637

DATE: 01/23/2001

TIME: 12:19:41

Input Set : A:\Cpg.pto

Output Set : N:\CRF3\01232001\1749637.raw

107 <210> SEQ ID NO: 2

109 <211> LENGTH: 37

111 <212> TYPE: PRT

113 <213> ORGANISM: *Comus gloriamaris*

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119 Met Lys Leu Thr Cys Met Met Ile Val Ala Val Leu Phe Leu Thr Ala

120 1 5 10 15

121 Trp Thr Thr Val Thr Ala Asp Asp Ser Gly Asn Gly Met Glu Ile Leu

122 20 25 30

123 Phe Pro Phe Val Gly Phe Glu Met Glu Asn Leu Glu Val Ser Asn Arg

124 35 40 45

125 Val Lys Phe Cys Arg Lys Glu Gly Gln Leu Cys Asp Pro Ile Phe Gln

126 50 55 60

127 Asn Cys Cys Arg Gly Trp Asn Cys Val Leu Phe Cys Val

128 65 70 75

130 <210> SEQ ID NO: 3

131 <211> LENGTH: 29

133 <212> TYPE: PRT

135 <213> ORGANISM: *Comus gloriamaris*

139 <214> FEATURE:

141 <221> NAME/KEY: SITE

143 <222> LOCATION: 3..(1)..(22)

145 <223> OTHER INFORMATION: Xaa at residues 3 and 13 may be pro or hydroxy-

Pro; Xaa at residue

146 1 5 10 15 20 25 30 35 40 45 50 55 60 65 70 75 80 85 90 95

147 100 105 110 115 120 125 130 135 140 145 150 155 160 165 170 175 180 185 190 195 200 205 210 215 220 225 230 235 240 245 250 255 260 265 270 275 280 285 290 295 300 305 310 315 320 325 330 335 340 345 350 355 360 365 370 375 380 385 390 395 400

148 400 SEQUENCE: 3

W--> 163 Val Lys Xaa Cys Arg Lys Xaa Gly Gln Leu Cys Asp Xaa Ile Phe Gln

164 1 5 10 15

W--> 166 Asn Cys Cys Arg Gly Xaa Asn Cys Val Leu Phe Cys Val

167 20 25

169 <210> SEQ ID NO: 4

171 <211> LENGTH: 29

173 <212> TYPE: PRT

175 <213> ORGANISM: *Comus gloriamaris*

179 <214> FEATURE:

181 <221> NAME/KEY: SITE

183 <222> LOCATION: (1)..(29)

185 <223> OTHER INFORMATION: Xaa at residues 3 and 13 may be pro or hydroxy-

Pro; Xaa at residue

186 1 5 10 15 20 25 30 35 40 45 50 55 60 65 70 75 80 85 90 95

187 100 105 110 115 120 125 130 135 140 145 150 155 160 165 170 175 180 185 190 195 200 205 210 215 220 225 230 235 240 245 250 255 260 265 270 275 280 285 290 295 300 305 310 315 320 325 330 335 340 345 350 355 360 365 370 375 380 385 390 395 400

188 400 SEQUENCE: 4

W--> 194 Val Lys Xaa Cys Arg Lys Xaa Gly Gln Leu Cys Asp Xaa Ile Xaa Gln

195 1 5 10 15

W--> 197 Asn Cys Cys Arg Gly Xaa Asn Cys Val Leu Phe Cys Val

198 20 25

200 <210> SEQ ID NO: 5

202 <211> LENGTH: 29

204 <212> TYPE: PRT

RAW SEQUENCE LISTING

PATENT APPLICATION: US/09/749,637

DATE: 01/23/2001

TIME: 12:19:41

Input Set : A:\Cpg.pto

Output Set: N:\CRF3\01232001\I749637.raw

200 <213> ORGANISM: *Conus gloriamaris*

210 <220> FEATURE:

212 <221> NAME/KEY: SITE

214 <222> LOCATION: (1)..(29)

216 <223> OTHER INFORMATION: Xaa at residues 3 and 13 may be pro or hydroxy-

pro; Xaa at residu

217 e 7 may be Gln or gamma-carboxy-Gln; Xaa at residue 22 may be His

218 or homo-His; Xaa at residue 27 may be Tyr, 12'-O-Tyr, mono-metho-

219 -Tyr, di-metho-Tyr, O-sulpho-Tyr or O-phospho-Tyr

223 <400> SEQUENCE: 5

W--> 225 Val Lys Xaa Cys Arg Lys Xaa Gly Gln Leu Cys Asp Xaa Ile Phe Gln

226 1 5 10 15

W--> 228 Asn Cys Cys Arg Gly Xaa Asn Cys Val Leu Xaa Cys Val

229 20 25

231 <210> SEQ ID NO: 6

233 <211> LENGTH: 342

235 <212> TYPE: DNA

237 <213> ORGANISM: *Conus omaria*

241 <220> FEATURE:

243 <221> NAME/KEY: CDS

245 <222> LOCATION: (116)..(235)

247 <400> SEQUENCE: 6

249 gaagatgga cagctgaaag taccgtcga gattacggg atcgactca tcatcatcp 1

252 tcaatgtgc cctcatccc ttcattcat cactgcaga ctatataaa catcaatc 20

254 tctctctct tctgagttg acaga tcc atc aac tgg tgc cgt aga gaa gat 40

255 Ser Ile Arg Met Cys Arg Arg Glu Ala

256 1 5

258 caa att tgc gat ccc att ttt caa aac tgc tgc cat agc ttg ttt tgc 220

259 Gln Leu Cys Asp Pro Ile Phe Gln Asn Cys Cys His Gly Leu Phe Cys

260 10 15 20 25

262 gtt ttg ttc tgc gtc taaaactaac gtgactctt cctctccct ctagtacta 275

263 Val Leu Val Cys Val

264 30

266 taggcagccg ctctagaaag tccaaagctt cgtacccgtg catgcagcgt catagctctt 335

268 ctatagtgc accaaattc aattcaactgc ccttcgtttt aaacgctcgt gactgggaaa 395

270 accttgcgt tacczaatt aatcgccttg cagacatcc ccttttcgcc aactggccta 455

272 atagcgaaga ggcg gcaac atcgcctt ccaaacagtt gcc agcctg aatggcqaat 515

275 agaacgcgcc ctatagcgcg caattct 542

277 <210> SEQ ID NO: 7

279 <211> LENGTH: 30

281 <212> TYPE: PRT

283 <213> ORGANISM: *Conus omaria*

287 <400> SEQUENCE: 7

289 Ser Ile Arg Met Cys Arg Arg Glu Ala Gln Leu Cys Asp Pro Ile Phe

290 1 5 10 15

293 Gln Asn Cys Cys His Gly Leu Phe Cys Val Leu Val Cys Val

294 20 25 30

297 <210> SEQ ID NO: 8

299 <211> LENGTH: 27

301 <212> TYPE: PRT

RAW SEQUENCE LISTING

PATENT APPLICATION: US/09/749,637

DATE: 01/23/2001

TIME: 17:19:11

Input Set : A:\Cpg.pto

Output Set : N:\CRF3\01232001\I749637.raw

318 <213> ORGANISM: Conus omaria

319 <220> FEATURES:

320 <221> NAME/KEY: SITE

321 <222> LOCATION: (1)..(27)

323 <223> OTHER INFORMATION: Xaa at residue 1 is glu or glutamic-carboxy-glu. Xaa at residue 11 is

324

325 <400> SEQUENCE: 8

W--> 320 Met Cys Arg Arg Xaa Ala Gln Leu Cys Asp Xaa Ile Phe Gln Asn Cys

321 1 10

323 Cys His Glu Leu Phe Cys Val Leu Val Cys Val

324 20 25

326 <210> SEQ ID NO: 9

328 <211> LENGTH: 346

330 <212> TYPE: DNA

332 <213> ORGANISM: Conus textile

333 <220> FEATURES:

334 <221> NAME/KEY: CDS

340 <222> LOCATION: (27)..(315)

341 <400> SEQUENCE: 9

345 cgccttacct aagcatcac caag atg aaa ctg aca tgc atg ata atc att 31

346 Met Lys Leu Thr Cys Met Met Ile Val

347 5

349 cct ata ctg ttc tta acc ggc tgc acc ttc gtc cca att ggt ggc tgc 32

350 Ala Val Leu Phe Leu Thr Ala Trp Thr Phe Val Thr Ala Asp Asn Ser

351 10 15 20 25

353 aga aat gga atg gaa aat ctt ttc ccg aag gca ggt ctc gaa atg gaa 147

354 Arg Asn Glu Leu Glu Asn Leu Phe Pro Lys Ala Gly His Glu Met Glu

355 30 35 40

357 gac ctg ggc gac tct aaa cac aag cac cag gaa aga ccg gac acc ggc 195

358 Ser Leu Glu Asp Ser Lys His Arg His Glu Glu Arg Pro Asp Thr Gly

359 45 50 55

361 gac aaa gaa gaa atg ctg cta cag aga cac gtc aag ccg tgt cgt aaa 243

362 Asp Lys Glu Glu Met Leu Leu Gln Arg Gln Val Lys Pro Cys Arg Lys

363 60 65 70

365 gaa cat caa ctg tat gat ctg att ttt caa aac tgc tgc cgt agc tgc 291

366 Glu His Gln Leu Cys Asp Leu Ile Phe Gln Asn Cys Cys Arg Gly Trp

367 75 80 85

369 tat tgc att atc ctg tct tgc act tgaagaetac ctgatgtgtt ctactccat 345

370 Trp Cys Val Val Leu Ser Cys Thr

371 90 95

373 c 346

376 <210> SEQ ID NO: 10

378 <211> LENGTH: 97

380 <212> TYPE: PRP

382 <213> ORGANISM: Conus textile

386 <400> SEQUENCE: 10

388 Met Lys Leu Thr Cys Met Met Ile Val Ala Val Leu Phe Leu Thr Ala

389 1 5 10 15

392 Trp Thr Phe Val Thr Ala Asp Asp Ser Arg Asn Gly Met Glu Asn Leu

RAW SEQUENCE LISTING

PATENT APPLICATION: US/09/749,637

DATE: 01/23/2001

TIME: 12:19:41

Input Set : A:\Cpg.pto

Output Set : N:\CRF3\01232001\I749637.raw

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100      20      30      40      50
101 the pro thr ala gly tyr gln met gln asn leu thr asp ser his his
102      35      40      50
103 ala his gln gln arg pro asp thr gln asp lys glu gln met leu leu
104      10      35      50
105 gln arg gln val lys pro cys arg lys gln his gln leu cys asp leu
106      65      70      75      80
107 thr phe gln asn cys arg gly arg leu ala val val leu ser cys
108      60      70      75      80
109
110

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111 <210> SEQ ID NO: 11

112 <211> LENGTH: 81

113 <212> TYPE: PRO

114 <213> ORGANISM: Camus textile

115 <220> FEATURE:

116 <221> NAME/KEY: CDS

117 <222> LOCATION: (1)..(81)

118 <223> OTHER INFORMATION: Xaa at residue 1 may be Gln or pyrolysine. Xaa at residue 4 may be

119 pro or hydroxy-pro. Xaa at residue 23 may be thr or homo-thr. Xaa

120 at residue 24 may be thr, leu or tyr. mono-iodo-tyr or di-iodo-tyr

121 phosphorylation or 6-phosphorylation

122 <224> REMARK: 11

W--> 441 Xaa Val Lys Xaa Cys Arg Lys Xaa His Gln Leu Cys Asp Leu Ile Phe

```

123      5      10      15      20

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W--> 444 Gln Asn Cys Cys Arg Gly Xaa Xaa Cys Val Val Leu Ser Cys Thr

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124      20      25      30

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125 <210> SEQ ID NO: 12

126 <211> LENGTH: 165

127 <212> TYPE: DNA

128 <213> ORGANISM: Conus omaria

129 <220> FEATURE:

130 <221> NAME/KEY: CDS

131 <222> LOCATION: (1)..(234)

132 <400> SEQUENCE: 12

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133 atc aac cta acc agc ctg atg atc att gaa atg ctg tcc ttg acc ggc      48

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134 Met Lys Leu Thr Cys Leu Met Ile Val Ala Val Leu Ser Leu Thr Gly

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135      5      10      15

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136 tgg aca ttc gtc aag gct gat gac tct gaa aat gga ttg ggg aat ctt      96

```

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137 Trp Thr Phe Val Thr Ala Asp Asp Ser Gly Asn Gly Leu Gly Asn Leu

```

```

138      20      30

```

```

139 ttt tgg aat gaa cat cac gaa atg aag aac ccc gaa gcc tct aaa ttg      144

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140 Phe Ser Asn Ala His His Glu Met Lys Asn Pro Gln Ala Ser Lys Leu

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```

141      35      40      45

```

```

142 aac aag agg tgc gtt cca ccc gag gcc cct tgt aat tgg ctt aca caa      192

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```

143 Asn Lys Arg Cys Val Pro His Glu Gly Pro Cys Asn Trp Leu Thr Gln

```

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144      50      55      60

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145 aac tgc tgc aat ggt tat aat tgc atc att ttt ttc tgc cta ~      234

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146 Asn Cys Cys Ser Gly Tyr Asn Cys Ile Ile Phe Phe Cys Leu

```

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147      65      70      75

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Please Note:

Use of n and/or Xaa have been detected in the Sequence Listing. Please review the Sequence Listing to ensure that a corresponding explanation is presented in the <220> to <223> fields of each sequence which presents at least one n or Xaa.

VERIFICATION SUMMARY

PATENT APPLICATION: US/09/749,637

DATE: 11/13/2001

TIME: 12:11:42

Input Set : A:\Cpg.pto

Output Set : N:\CRF3\01232001\I749637.raw

L:83 H:270 C: Current Application Number differs. Replaced Current Application No

L:85 H:281 C: Current filing date differs. Replaced Current Filing Date

L:165 H:341 W: (46) "n" or "Xaa" used, for SEQ ID# 3

L:166 H:341 W: (46) "n" or "Xaa" used, for SEQ ID# 3

L:194 M:341 W: (46) "n" or "Xaa" used, for SEQ ID# 4

L:195 M:341 W: (46) "n" or "Xaa" used, for SEQ ID# 4

L:225 H:341 W: (46) "n" or "Xaa" used, for SEQ ID# 5

L:226 M:341 W: (46) "n" or "Xaa" used, for SEQ ID# 5

L:229 H:341 W: (46) "n" or "Xaa" used, for SEQ ID# 3

L:341 H:341 W: (46) "n" or "Xaa" used, for SEQ ID# 17

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L:198 L:341 W: (46) "n" or "Xaa" used, for SEQ ID# 23

L:199 M:341 W: (46) "n" or "Xaa" used, for SEQ ID# 23

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L:217 M:341 W: (46) "n" or "Xaa" used, for SEQ ID# 26

L:1058 H:141 W: (46) "n" or "Xaa" used, for SEQ ID#12

L:106 H:141 W: (46) "n" or "Xaa" used, for SEQ ID#12

L:1068 H:141 W: (46) "n" or "Xaa" used, for SEQ ID#120

L:1071 H:141 W: (46) "n" or "Xaa" used, for SEQ ID#128

L:1172 H:341 W: (46) "n" or "Xaa" used, for SEQ ID#31

L:1175 H:341 W: (46) "n" or "Xaa" used, for SEQ ID#31

L:1277 H:341 W: (46) "n" or "Xaa" used, for SEQ ID#34

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L:1600 H:341 W: (46) "n" or "Xaa" used, for SEQ ID#43

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L:1920 H:341 W: (46) "n" or "Xaa" used, for SEQ ID#52

L:1946 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#53

L:1976 H:341 W: (46) "n" or "Xaa" used, for SEQ ID#54

L:1979 H:341 W: (46) "n" or "Xaa" used, for SEQ ID#54

L:2079 H:341 W: (46) "n" or "Xaa" used, for SEQ ID#57

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L:2285 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#63

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VERIFICATION SUMMARY

PATENT APPLICATION: US/09/749,637

DATE: 01/23/2001

TIME: 12:19:42

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Output Set: N:\CRF3\01232001\I749637.raw

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L12493 H:341 W: (46) "n" or "Xaa" used. for SEQ ID#168
L12496 H:341 W: (46) "n" or "Xaa" used. for SEQ ID#169

OIIPE

RAW SEQUENCE LISTING

PATENT APPLICATION: US,09/749,637

DATE: 01/16/2001

TIME: 11:53:59

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Output Set : N:\CRF3\01162001\I749637.raw

1 <110> APPLICANT: University of Utah Research Foundation
 2 <110> Cognetix, Inc.
 3 <110> Glicer, Baldomero H.
 4 <110> Cartier, G. Edward
 5 <110> Watkins, Martin
 6 <110> Hill, David R.
 7 <110> Netherland, J. Michael
 8 <110> Layer, Richard L.
 9 <110> Jones, Robert E.
 10 <110> TITLE OF INVENTION: O-Supranuclear Core-2 In Peptide
 11 <110> FILE REFERENCE: 2314-227
 C--> 17 <140> CURRENT APPLICATION NUMBER: US,09/749,637
 C--> 17 <141> CURRENT FILING DATE: 2000-12-28
 17 <150> PRIOR APPLICATION NUMBER: US 60/213,112
 18 <151> PRIOR FILING DATE: 2000-10-27
 20 <150> PRIOR APPLICATION NUMBER: US60/219,440
 21 <151> PRIOR FILING DATE: 2000-07-20
 23 <150> PRIOR APPLICATION NUMBER: US 60/211,263
 24 <151> PRIOR FILING DATE: 2000-06-26
 26 <150> PRIOR APPLICATION NUMBER: US 60/173,754
 27 <151> PRIOR FILING DATE: 1999-12-30
 29 <160> NUMBER OF SEQ ID NOS: 409
 31 <170> SOFTWARE: Patent, version 3.0

Does Not Comply
Corrected Diskette Needed
PP 1, 2

These amino acids
were not found.

PROBSED SEQUENCES

959 <210> SEQ ID NO: 24
 960 <211> LENGTH: 27
 961 <212> TYPE: PRT
 962 <213> ORGANISM: Corus ammiralis
 964 <220> FEATURE:
 965 <221> NAME/KEY: SITE
 966 <222> LOCATION: (1)..(27)
 967 <223> OTHER INFORMATION: Xaa at residue 1 may be Trp or bromo-Trp; Xaa at
 residues 7 and 1

968 <223> 8 may be Glu or gamma-carboxy-Glu; Xaa at residue 20 may be Tyr
 969 <223> 125-I-Tyr, mono-iodo-Tyr, di-iodo-Tyr, O-sulpho-Tyr or O-phospho-

E--> 970 <223> Tyr
 971 Tyr
 973 <400> SEQUENCE: 11
 E--> 975 Xaa Cys Lys Glu Ser Gly Xaa Met Cys Asn Leu Leu Asp Gln Asn Cys
 976 1 5 10 15
 W--> 978 Cys Xaa Gly Xaa Cys Ile Val Leu Val Cys Thr
 979 20 25

8367 <210> SEQ ID NO: 313

8368 <211> LENGTH: 26

8369 <212> TYPE: PRT

move to here or place <223> before each
 line of the
 explanation.
 When a lone amino acid is placed
 like this and does not have a <223>
 in front of it, the computer program
 picks it up as the first amino in

RAW SEQUENCE LISTING

PATENT APPLICATION: US/09/749,637

DATE: 01/16/2001

TIME: 11:53:40

Input Set : A:\2314-227.ST25.txt

Output Set : N:\CRF3\01162001\1749637.raw

8370 <213> ORGANISM: Canis familiaris

8371 <219> CHARACTER:

8372 <221> NAME/KEY: S111

8373 <222> LOCATION: (1)..(26)

8374 <223> OTHER INFORMATION: Xaa at residues 3 and 7 may be Trp or bromo-Trp;

Xaa at residues

8375 <223> 1 and 17 may be Pro or hydroxy-Pro; Xaa at residue 6 may be Tyr;

8376 <223> 18 may be Tyr, mono-iodo-Tyr, di-iodo-Tyr, O-sulpho-Tyr or O-phospho-

E--> 8378 <223> Tyr

8379

8380 <400> SEQUENCE: A13

E--> 8383 Asp Cys Xaa Xaa Gln Xaa Xaa Phe Cys Gly Leu Gln Arg Gly Cys Cys

8384 1 5 10 15

W--> 8386 Xaa Gly Thr Thr Cys Phe Phe Leu Cys Phe

8387

20

25

*move to here or place <223> before each**line of
the explanatory**Same error, refer
to p.1.*

VERIFICATION SUMMARY

PATENT APPLICATION: US/09/749,637

DATE: 01/16/2001

TIME: 11:33:11

Input Set : A:\2314-227.ST25.txt

Output Set : N:\CRF3\01162001\I749637.raw

L.117 H:250 F: Current Application Number differs. Replaced Current Application No.
 L.117 H:250 F: Current Filing Date differs. Replaced Current Filing Date
 L.118 H:341 W: (46) "n" or "Xaa" used, for SEQ ID# 3
 L.119 H:341 W: (46) "n" or "Xaa" used, for SEQ ID# 3
 L.120 H:341 W: (46) "n" or "Xaa" used, for SEQ ID# 4
 L.121 H:341 W: (46) "n" or "Xaa" used, for SEQ ID# 4
 L.122 H:341 W: (46) "n" or "Xaa" used, for SEQ ID# 5
 L.123 H:341 W: (46) "n" or "Xaa" used, for SEQ ID# 5
 L.124 H:341 W: (46) "n" or "Xaa" used, for SEQ ID# 6
 L.125 H:341 W: (46) "n" or "Xaa" used, for SEQ ID# 6
 L.126 H:341 W: (46) "n" or "Xaa" used, for SEQ ID# 11
 L.127 H:341 W: (46) "n" or "Xaa" used, for SEQ ID# 11
 L.128 H:341 W: (46) "n" or "Xaa" used, for SEQ ID# 11
 L.129 H:341 W: (46) "n" or "Xaa" used, for SEQ ID# 14
 L.130 H:341 W: (46) "n" or "Xaa" used, for SEQ ID# 14
 L.131 H:341 W: (46) "n" or "Xaa" used, for SEQ ID# 17
 L.132 H:341 W: (46) "n" or "Xaa" used, for SEQ ID# 17
 L.133 H:341 W: (46) "n" or "Xaa" used, for SEQ ID# 20
 L.134 H:341 W: (46) "n" or "Xaa" used, for SEQ ID# 20
 L.135 H:341 W: (46) "n" or "Xaa" used, for SEQ ID# 23
 L.136 H:341 W: (46) "n" or "Xaa" used, for SEQ ID# 23
 L.137 H:341 W: (46) "n" or "Xaa" used, for SEQ ID# 24
 L.138 H:341 W: (46) "n" or "Xaa" used, for SEQ ID# 24
 L.139 H:341 W: (46) "n" or "Xaa" used, for SEQ ID# 27
 L.140 H:341 W: (46) "n" or "Xaa" used, for SEQ ID# 27
 L.141 H:341 W: (46) "n" or "Xaa" used, for SEQ ID# 28
 L.142 H:341 W: (46) "n" or "Xaa" used, for SEQ ID# 28
 L.143 H:341 W: (46) "n" or "Xaa" used, for SEQ ID# 31
 L.144 H:341 W: (46) "n" or "Xaa" used, for SEQ ID# 31
 L.145 H:341 W: (46) "n" or "Xaa" used, for SEQ ID# 31
 L.146 H:341 W: (46) "n" or "Xaa" used, for SEQ ID# 34
 L.147 H:341 W: (46) "n" or "Xaa" used, for SEQ ID# 34
 L.148 H:341 W: (46) "n" or "Xaa" used, for SEQ ID# 37
 L.149 H:341 W: (46) "n" or "Xaa" used, for SEQ ID# 37
 L.150 H:341 W: (46) "n" or "Xaa" used, for SEQ ID# 37
 L.151 H:341 W: (46) "n" or "Xaa" used, for SEQ ID# 40
 L.152 H:341 W: (46) "n" or "Xaa" used, for SEQ ID# 40
 L.153 H:341 W: (46) "n" or "Xaa" used, for SEQ ID# 40
 L.154 H:341 W: (46) "n" or "Xaa" used, for SEQ ID# 43
 L.155 H:341 W: (46) "n" or "Xaa" used, for SEQ ID# 43
 L.156 H:341 W: (46) "n" or "Xaa" used, for SEQ ID# 46
 L.157 H:341 W: (46) "n" or "Xaa" used, for SEQ ID# 46
 L.158 H:341 W: (46) "n" or "Xaa" used, for SEQ ID# 49
 L.159 H:341 W: (46) "n" or "Xaa" used, for SEQ ID# 49
 L.160 H:341 W: (46) "n" or "Xaa" used, for SEQ ID# 52
 L.161 H:341 W: (46) "n" or "Xaa" used, for SEQ ID# 52
 L.162 H:341 W: (46) "n" or "Xaa" used, for SEQ ID# 52
 L.163 H:341 W: (46) "n" or "Xaa" used, for SEQ ID# 53
 L.164 H:341 W: (46) "n" or "Xaa" used, for SEQ ID# 53
 L.165 H:341 W: (46) "n" or "Xaa" used, for SEQ ID# 54
 L.166 H:341 W: (46) "n" or "Xaa" used, for SEQ ID# 54
 L.167 H:341 W: (46) "n" or "Xaa" used, for SEQ ID# 57
 L.168 H:341 W: (46) "n" or "Xaa" used, for SEQ ID# 57
 L.169 H:341 W: (46) "n" or "Xaa" used, for SEQ ID# 57
 L.170 H:341 W: (46) "n" or "Xaa" used, for SEQ ID# 57
 L.171 H:341 W: (46) "n" or "Xaa" used, for SEQ ID# 60
 L.172 H:341 W: (46) "n" or "Xaa" used, for SEQ ID# 60
 L.173 H:341 W: (46) "n" or "Xaa" used, for SEQ ID# 60
 L.174 H:341 W: (46) "n" or "Xaa" used, for SEQ ID# 63
 L.175 H:341 W: (46) "n" or "Xaa" used, for SEQ ID# 63

VERIFICATION SUMMARY

PATENT APPLICATION: US/09/749,637

Date: 1/16/2001

Time: 11:54:11

Input Set : A:\2314-227.ST25.txt

Output Set : N:\CRF3\01162001\I749637.raw

1:176: M:341 W: (46) "L" or "Xaa" used. for Seq ID#100
 1:176: H:341 W: (46) "L" or "Xaa" used. for Seq ID#100
 1:184: E:341 W: (46) "L" or "Xaa" used. for Seq ID#100
 1:192: E:341 W: (46) "L" or "Xaa" used. for Seq ID#100
 1:192: M:341 W: (46) "L" or "Xaa" used. for Seq ID#100
 1:3878: H:252 E: No. of Seq. differs. including Input:26 E: 111. Seq. no.